

The mouse Gene Expression Database (GXD): 2011 update

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Received September 15, 2010; Accepted October 1, 2010

ABSTRACT

The Gene Expression Database (GXD) is a community resource of mouse developmental expression information. GXD integrates different types of expression data at the transcript and protein level and captures expression information from many different mouse strains and mutants. GXD places these data in the larger biological context through integration with other Mouse Genome Informatics (MGI) resources and interconnections with many other databases. Web-based query forms support simple or complex searches that take advantage of all these integrated data. The data in GXD are obtained from the literature, from individual laboratories, and from large-scale data providers. All data are annotated and reviewed by GXD curators. Since the last report, the GXD data content has increased significantly, the interface and data displays have been improved, new querying capabilities were implemented, and links to other expression resources were added. GXD is available through the MGI web site (www.informatics.jax.org), or directly at www.informatics.jax.org/expression.shtml.

INTRODUCTION

As a primary mammalian model of human disease, the mouse is used extensively for expression studies to determine the role of genes that function in molecular pathways during developmental and disease processes. With a focus on endogenous gene expression during development, the Gene Expression Database (GXD) collects data from the scientific literature, from individual laboratories, and from large-scale data providers. It makes these data readily available to the research community in a highly curated and integrated format that allows for a large variety of

database queries. GXD captures a broad spectrum of assay types, including RNA *in situ* hybridization, immunohistochemistry, knock-in reporter assays, northern blot, western blot, RT-PCR, RNase protection and S1 nuclease assays. It covers all developmental stages and tissues and includes data from many different mouse strains and mutants, giving researchers a tool to examine the effects of mutations on gene expression. GXD forms an important and integral component of the larger Mouse Genome Informatics (MGI) resource. Therefore, the expression data are fully integrated with mouse genetic, sequence, functional and phenotypic information (1–4). MGI maintains further links to many other resources such as GenBank, gene model resources, Entrez Gene, UniProt, InterPro, Online Mendelian Inheritance in Man (OMIM) and the International Knockout Mouse Consortium (IKMC) among others (5–14). This robust integration puts the expression data in GXD into a much larger biological and analytical context.

Other databases that store mouse expression information have been developed in recent years. They store data from one or two specific assay types and/or focus on specific developmental stages; they are often dedicated to specific data generation projects (15–22). As will be evident from this article, GXD is working with those resources that are complementary to GXD, adding value through data integration and the implementation of new interconnections. Due to its broad scope, its extensive data curation and integration efforts, and the resulting querying capabilities, GXD continues to provide a unique resource to the biomedical research community. GXD is updated daily. GXD and its query interfaces have been described earlier (23–27). Here, we report on our recent progress in terms of data acquisition, and on the implementation of new query and display features.

NEW GXD HOMEPAGE

To present the objectives of GXD more clearly and to make the database more intuitive to use, we redesigned

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the GXD homepage (www.informatics.jax.org/expression.shtml). The new layout provides clear access to the various query forms, with short descriptors for each form. The ‘Frequently Asked Questions (FAQ’s)’ section provides links to brief on-line tutorials demonstrating how one can search for different types of data in GXD. The ‘GXD Includes’ section provides information about the current data content in GXD, such as the number of genes with annotated expression data, the number of expression results and the number of images in the database. The ‘Gene Expression News’ section informs users when new features, capabilities and data sets become available. A series of tabs at the bottom of the home page provides access to help documentation and data policies, information about GXD and its collaborators, and links to guidelines and tools that help researchers to submit data electronically. That GXD is also an integral component of MGI is made transparent through the use of a central ‘Quick Search’ (see below), a common navigation bar and common drop-menus and tab choices that direct users to various data sets, search forms, tools and other resources. Large icons on the MGI homepage provide visual cues to the various core areas, including expression (GXD).

DATA CONTENT

Literature Summary

In the Literature Summary, GXD provides users with a way to quickly determine what mouse developmental expression data are available in the scientific literature. The staff of GXD searches the scientific literature for publications that present endogenous gene expression experiments during mouse development. In a first annotation step for each publication, the genes analyzed, the ages of mice used in the experiment, and the type of experimental assay performed for each gene are recorded and entered into the database. These data are easily searched using the ‘Gene Expression Literature Query Form’. These queries can also include citation (author, journal, year) and abstract information. However, this tool takes users further than a Pubmed search because the data in the GXD Literature Summary are based on the curation of the full-text of the paper, including supplemental information, and annotations are standardized with regard to gene, age and assay type information. The Literature Summary is comprehensive and up-to-date. It includes all journal articles containing expression data during mouse development from 1993 to the present and all articles from major developmental journals since 1990. Currently, the GXD Literature Summary has 108 604 records covering data for 13 619 genes from 17 521 references.

Gene expression data

Beyond summaries, GXD also provides detailed records of experimental expression results. GXD assay records contain the authors’ description of the tissue pattern and strength of expression, translated into standard terminology (see below), the probe or antibody information

available, as well as the specimen age, genetic background and preparation (Figure 1). The expression information is recorded in standardized formats by making extensive use of controlled vocabularies and ontologies, thus enabling data integration and complex querying capabilities. To capture the author’s descriptions of where expression was or was not detected, GXD uses a dictionary of anatomical terms that lists the anatomical structures for each developmental stage in a hierarchical fashion. In this way, expression data from assays with differing spatial resolution can be recorded in a consistent manner, and the expression pattern information becomes accessible to hierarchical searches. The developmental part of the anatomical dictionary was established by our collaborators from the EMAGE project (28) and is being extended and refined jointly; the postnatal part was developed by the GXD project (29).

In addition to curating expression data from the literature on a daily basis, GXD continues to work with individual laboratories and large-scale data providers to bring their data sets into the database. Recently acquired large data sets include: an RT-PCR screen of more than 800 genes in pre-implantation embryos (30); an RNA *in situ* hybridization screen of 293 genes at E7.5 (31); an RNA *in situ* hybridization screen of more than 200 genes in the developing hypothalamus (32); a set of knock-in lacZ reporter studies from Deltagen Inc.; an RNA *in situ* hybridization data for 745 genes at E14.5 from GenePaint (22) and a data set from the Eurexpress project (www.eurexpress.org), covering RNA *in situ* hybridization data for 6409 genes at E14.5. In all these cases, GXD staff worked with the data providers to bring the data into a standardized format that could be incorporated into GXD and to resolve issues pertinent to nomenclature, incompleteness, and referential integrity. Upon computational and manual review, the data were bulk-loaded into GXD.

Due to all these efforts, the content of expression data in GXD has increased tremendously in recent years. The database currently holds almost 930 000 expression results from 45 305 expression assays for 12 139 genes. This includes expression data from 1503 different mouse mutants.

ACCESSING GXD’S EXPRESSION DATA

An effective way to access the expression information for a single gene is to look up the ‘Expression’ section on the corresponding gene detail page (Figure 2). Gene detail pages can be easily found via the ‘Quick Search’ tool placed at the top of all MGI pages. Because these pages are a central hub within MGI, we have improved the layout of the expression section, implemented a new images summary (see below), and added new gene-specific links to external expression resources, including the Allen Brain Atlas (18), GENSAT (17), GEO (33) and ArrayExpress (34). Future work will include the implementation of links to model organisms from other species that hold developmental expression data, such as ZFIN (35), GEISHA (36), Xenbase (37) and Flybase (38).

Gene Expression Data

Your Input Welcome

Reference: J:67152 Perea-Gomez A, Development 2001 Mar;128(5):753-65
Assay type: RNA in situ
MGI Accession ID: MGI:1930293
Gene symbol: *Wnt3*
Gene name: wingless-related MMTV integration site 3
Modification date: 2/1/2006

Probe: *Wnt3* probe2
Visualized with: Autoradiography

Specimens Used

	7. B	7. B'
Genetic Background	Not Specified	involves: 129S1/Sv * 129X1/SvJ = CD-1
Mutant Allele(s)	<i>Otx2^{tm1Sls}/Otx2^{tm1Sls}</i>	
Age	E 6.5 (a)	E 6.5 (a)
Sex	Not Specified	Not Specified
Type	section	section
Fixation	4% Paraformaldehyde	4% Paraformaldehyde
Embedding	Cryosection	Cryosection

Notes:
(a) Age of embryo at noon of plug day not specified in reference.

Results: 7. B (embryonic day 6.5)

Structure	Level	Pattern	Note	Image
TSG: embryo; ectoderm	Present	Regionally restricted	(a)	Figure 7B
TSG: embryo; mesoderm	Present	Regionally restricted	(a)	Figure 7B

Notes:
(a) Expression is detected in the posterior epiblast and mesodermal cells.

Results: 7. B' (embryonic day 6.5; involves: 129S1/Sv * 129X1/SvJ = CD-1; *Otx2^{tm1Sls}/Otx2^{tm1Sls}*)

Structure	Level	Pattern	Note	Image
TSG: embryo; ectoderm	Present	Regionally restricted	(a)	Figure 7B'
TSG: embryo; mesoderm	Present	Regionally restricted	(a)	Figure 7B'

Notes:
(a) Expression is detected in the entire proximal epiblast and mesodermal cells.

Images

Query Results -- Details

Reference: J:67152 Perea-Gomez A, Development 2001 Mar;128(5):753-65
Figure: 7
MGI Accession ID: MGI:1930293
Assays that refer to this image: See below

Wnt3 Otx2 Bmp4 Nodal

WT

Otx2^{-/-}

Copyright: This image is from Perea-Gomez A, Development 2001;128(5):753-765, and is displayed with the permission of The Company of Biologists Limited who owns the Copyright.
Assays that refer to this image:

Label	Assay & Result Details (Gene Symbol)	Spatial Mapping
A	MGI:1930275 (<i>Otx2</i>)	
A'	MGI:1930275 (<i>Otx2</i>)	
B	MGI:1930293 (<i>Wnt3</i>)	EMAGE:542
B'	MGI:1930293 (<i>Wnt3</i>)	
C	MGI:1930275 (<i>Otx2</i>)	
C'	MGI:1930275 (<i>Otx2</i>)	
D	MGI:1930292 (<i>Bmp4</i>)	EMAGE:132
D'	MGI:1930292 (<i>Bmp4</i>)	
E	MGI:1930294 (<i>Nodal</i>)	EMAGE:685
E'	MGI:1930294 (<i>Nodal</i>)	
F	MGI:1930294 (<i>Nodal</i>)	
F'	MGI:1930294 (<i>Nodal</i>)	

Figure 1. Detailed gene expression data annotations and images. An assay record for *Wnt3* (left) and the accompanying image page (right). The assay page includes: publication reference, gene symbol and name, link to probe/antibody information, specimen details, and the expression annotations for each specimen. Annotations include the anatomical structures in which expression was analyzed and the strength of expression in these structures. They refer to corresponding figures in publications or electronic submissions. Whenever possible, the images are included in GXD and assay records link to those images. The image page includes: publication reference, figure label, the figure, a caption and a copyright statement of the journal or the submitter. Figures from publications often include several image panes analyzing different genes and/or genetic backgrounds. The table below the figure indicates each pane and its corresponding assay record with links to that record, as illustrated by the red circle. For images that have been spatially mapped onto the Edinburgh Mouse Atlas, the table provides a link to the corresponding entry in EMAGE.

The query forms available from the GXD homepage, as well as from the MGI search menu, provide more direct and more powerful means to access expression data. The ‘Gene Expression Data Query Form’ includes search fields that allow users to specify genes, tissue, developmental stage and expression assay type, as well as genome coordinates and gene ontology (GO) terms. Users can search for instances where expression was detected or not detected, as well as for expression data in specific mutants. Thus, this form enables both simple and very complex queries. The ‘Expanded Expression Data Query Form’ allows users to find genes that are expressed in one set of tissues and/or developmental stages, but not in others. The ‘Mouse Anatomical Dictionary Browser’ lets users search or browse for specific anatomical structures and the expression data associated with them. The ‘EMAGE Anatomical Section Browser’ uses the Edinburgh Mouse Atlas as a starting point for identifying anatomical structures and expression data associated with them in GXD and EMAGE.

Searches from the query form as well as the links from the expression section on the gene detail page lead to data summaries listing all assays/results that match the user’s criteria. We have improved the layout of these summaries.

They now feature clear ‘data’ links that take the user to the detailed entries, as well as camera icons that indicate if the detailed entries contain links to primary image data (Figures 1 and 2).

Other avenues in MGI can be used to access the expression data as well. We have increased the utility of the Quick Search tool by adding the capability to return expression results if the user enters anatomical terms. We have expanded the Batch Query tool so that it can return GXD expression data for a list of genes or sequence IDs, as well as export expression data as Excel worksheets or tab-delimited text files (Figure 3).

In addition to implementing gene specific links to the microarray gene expression data at GEO and ArrayExpress, we have, in collaboration with the Mouse Genome Database (MGD) project, developed new tools that allow researchers to combine microarray expression results with data in MGI. Specifically, we have begun to provide up-to-date mappings between microarray probe sets and genes so that users can query MGI by microarray probe set ID to retrieve the associated gene, or download reports of all probe set-to-gene mappings. We have also enhanced the MGI Batch Query tool so that it accepts sets of microarray probe IDs. Therefore, one can easily extract

Pax6
paired box gene 6
MGI:97490

Expression

Literature Summary: (644 records)
Data Summary: Assays (109) Results (1018) Tissues (438) Images (367)
Theiler Stages: 2,3,4,5,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,28

Assay Type	Assays	Results
Western blot	2	5
RT-PCR	11	63
RNA in situ	66	764
Immunohistochemistry	24	120
Northern blot	4	35
RNAse protection	2	31

cDNA source data(36)
External Resources: [Allen Brain Atlas](#) [GENSAT](#) [GEO](#) [ArrayExpress](#)

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Gene Expression Data Query Results -- Summary

109 matching assays displayed

Result Details	Gene	Assay Type	Reference
data (MGI:4196019)	Pax6	RT-PCR	J:98077 Alfano G, Hum Mol Genet 2004
data (MGI:2178971)	Pax6	Immunohistochemistry	J:76132 Aruga J, Dev Biol 2002 Apr
data (MGI:2158422)	Pax6	Immunohistochemistry	J:73074 Bai CB, Development 2001
data (MGI:2576442)	Pax6	Immunohistochemistry	J:79392 Bai CB, Development 2002
data (MGI:2386631)	Pax6	Immunohistochemistry	J:78684 Baumer N, Development 2002
data (MGI:2686871)	Pax6	RT-PCR	J:54475 Beimenes S, Mol Endocrinol 2002
data (MGI:2686872)	Pax6	RT-PCR	J:54475 Beimenes S, Mol Endocrinol 2002
data (MGI:2153747)	Pax6	RNA in situ	J:71879 Bernier G, Development 2001
data (MGI:2153750)	Pax6	RNA in situ	J:71879 Bernier G, Development 2001
data (MGI:3589693)	Pax6	RNA in situ	J:93300 Blackshaw S, PLoS Biol 2004
data (MGI:3039932)	Pax6	RNA in situ	J:76815 Broccoli V, Mech Dev 2002
data (MGI:2449580)	Pax6	RNA in situ	J:45299 Caric D, Development 1997
data (MGI:3054848)	Pax6	RNA in situ	J:89228 Chen MH, Genes Dev 2004
data (MGI:2159267)	Pax6	RT-PCR	J:74329 Clark SW, J Biol Chem 2002
data (MGI:2153172)	Pax6	Immunohistochemistry	J:65298 Corbin JG, Development 2001
data (MGI:2153173)	Pax6	RNA in situ	J:65298 Corbin JG, Development 2001
data (MGI:2676225)	Pax6	Immunohistochemistry	J:85473 Corbin JG, Development 2001
data (MGI:3711940)	Pax6	RNA in situ	J:47920 Dattani MT, Nat Genet 1998
data (MGI:1279437)	Pax6	Immunohistochemistry	J:49164 Ding Q, Development 1998
data (MGI:1322194)	Pax6	RNA in situ	J:49164 Ding Q, Development 1998
data (MGI:2179437)	Pax6	Immunohistochemistry	J:73778 Estivill-Torrus G, Development 1998
data (MGI:2157343)	Pax6	Immunohistochemistry	J:73525 Faber SC, Development 2001
data (MGI:2158660)	Pax6	Western blot	J:73625 Favari J, Genetics 2001 Dec
data (MGI:1338530)	Pax6	RNA in situ	J:41809 Filosa S, Development 1997
data (MGI:3777771)	Pax6	RNA in situ	J:122557 Furushima K, Dev Biol 2001
data (MGI:1340189)	Pax6	RNA in situ	J:51570 Furuta Y, Genes Dev 1998
data (MGI:1928094)	Pax6	Immunohistochemistry	J:65599 Gaufre GO, Development 2000
data (MGI:1342072)	Pax6	Immunohistochemistry	J:56424 Goodrich LV, Dev Biol 1999
data (MGI:3507622)	Pax6	RNA in situ	J:91257 Gray PA, Science 2004 Dec
data (MGI:3508832)	Pax6	RNA in situ	J:91257 Gray PA, Science 2004 Dec
data (MGI:3511921)	Pax6	RNA in situ	J:91257 Gray PA, Science 2004 Dec
data (MGI:3529950)	Pax6	RNA in situ	J:47200 Greene ND, Mech Dev 1998
data (MGI:3530041)	Pax6	RNA in situ	J:47700 Greene ND, Mech Dev 1998

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Gene Expression Data Images Query by Gene -- Summary

Symbol Name ID paired box gene 6 MGI:97490

367 expression image(s) for Pax6 in 151 figure(s)
Showing figure(s) 1 to 10 Show 10 figures per page Next Last

Click on thumbnail or figure label to view full size image with links to expression annotations.

Reference: J:149737 Sansom SN, et al. The level of the transcription factor Pax6 is essential for controlling the balance between neural stem cell self-renewal and neurogenesis. PLoS Genet 2009 Jun;5(6):e100511

Figure 4
B Wild type : Immunohistochemistry

Reference: J:149737 Sansom SN, et al. The level of the transcription factor Pax6 is essential for controlling the balance between neural stem cell self-renewal and neurogenesis. PLoS Genet 2009 Jun;5(6):e100511

Figure 9
A Pax6, A Pax6/Neurogen2/Hes1 : Immunohistochemistry

Reference: J:139177 Hoffman BG, et al. Identification of transcripts with enriched expression in the developing and adult pancreas. Genome Biol 2008;9(6):R99

Figure Embryo_C1193_1_3C
RNA in situ

Reference: J:122557 Furushima K, et al. Mouse homologues of Shisa antagonistic to Wnt and Fgf signalings. Dev Biol 2007 Jun 15;306(2):480-92

Figure 6
J : RNA in situ

Figure 2. Expression data summaries. Gene detail pages in MGI feature an ‘Expression’ section that summarizes the GXD data available for the gene from a variety of perspectives: literature content, assays, results, tissues, images, developmental stages and assay types (top). Links to the respective summaries provide quick access to the actual data. Gene-specific links to the Allen Brain Atlas, GENSAT, GEO and ArrayExpress are provided as well. Assay summaries (bottom left) link to the detailed assay records ('data') and show camera icons to indicate assay records with image links. The image summary (bottom right) lists thumbnails of all expression images for a gene and links to the full-size image page.

information about genes for sets of microarray probes, including RNA *in situ* hybridization, immunohistochemistry and RT-PCR expression data, GO annotations, mouse phenotype data and disease data associated with the orthologous human genes. Query results are available as web page displays and can also be downloaded as

tab-delimited text and thus fed into other analysis tools (Figure 3).

GXD is implemented in the Sybase relational database management system. If a user’s queries are beyond the capabilities of our web-based forms, custom SQLs or direct SQL access can be requested via our user support

MGI Batch Query

Download gene/marker data for a batch of IDs or symbols. Results may be in web or tab-delimited format.

Input

Type: Affy 1.0 ST
Source: Enter Text | Upload File

ID/Symbols List:
10413243
10576911
10434925
10419356
10474312
10528872

*tab, space, and newline separated IDs.

Output

Gene Attributes:
 Nomenclature Genome Location Ensembl ID Entrez Gene ID VEGA ID

Additional Information:
 Gene Ontology (GO) Mammalian Phenotype (MP) Human Disease (OMIM) Alleles
 Gene Expression RefSNP ID GenBank/RefSeq ID UniProt ID
 None

Format: Web (10,000 row maximum) Tab-delimited Text (200,000 row maximum) Excel (200,000 row maximum)

Search | Reset

Figure 3. MGI Batch Query. Batch queries can return GXD expression data for a list of genes or sequence IDs. Affymetrix probe set IDs are also accepted as input. Because MGI maintains up-to-date mappings of microarray probe set-to-gene associations, users can easily obtain MGI information about genes for specific sets of microarray probes, such as expression, phenotypic or, as shown, human disease data. Query results are available as web page displays, as tab-delimited text files or as Excel files. The Batch Query tool can be accessed at <http://www.informatics.jax.org/javawi2/servlet/WIFetch?page=batchQF>.

group. In addition, a number of reports for specific data sets, such as a list of all the genes that have annotated expression data, are produced nightly and are available on the website.

IMAGES IN GXD

Whenever possible, the standardized text annotations of expression data are linked to the primary image data. There are currently more than 191 099 images in GXD. Images are accessible via the ‘Expression’ section on the gene detail page. The recently implemented image summary displays all the images for the corresponding gene as thumbnails (Figure 2). These thumbnails are then linked to the full-sized image page. However, users are not restricted to selecting images on a per gene basis. Because all images are extensively indexed via the text annotations, they are accessible through the different search parameters that GXD provides on its query forms. Query result summaries display camera icons for entries that contain image data. The detailed entries then link directly to primary images obtained from the publication or from the data provider.

As part of our longstanding collaboration with the EMAGE project, which aims to ultimately create a resource that fully combines standardized text based and graphical means for storing and querying expression data (39), GXD makes all its *in situ* images and pertinent text annotations available to EMAGE. EMAGE then provides 3D spatial mappings of selected *in situ* data. The image detail pages in GXD now link to entries in EMAGE that

show the image spatially mapped onto the 3D atlas (Figure 1).

Large-scale *in situ* hybridization projects have contributed both data and images to GXD. Two of them, GenePaint and Eurexpress, provide additional images and features for image viewing at their web sites. Therefore, GXD has added links from the image page at GXD to the corresponding image entries at GenePaint and Eurexpress. These successful collaborations illustrate how community resources such as GXD and project-oriented resources such as GenePaint and Eurexpress can work with each other. GXD adds value to the incorporated data by integrating them with other expression data and with all the other types of data stored in MGI and by providing unique and powerful querying capabilities. Further, GXD maintains the data and keeps them current with regard to nomenclature and data connections. Conversely, the GenePaint and Eurexpress data significantly enrich GXD, and the links implemented to GenePaint and Eurexpress provide additional utility such as high-resolution images and zoom functions.

DIRECT ELECTRONIC CONTRIBUTION OF DATA

Users can contribute data directly to GXD. Guidelines for electronic submission are available on the web site, and a GXD staff member is available to assist. GXD has developed an Excel-based program, the Gene Expression Notebook (GEN), which provides users with a template for submitting expression data to GXD. A variety of other formats are also accepted. The GEN is also useful in the lab for maintaining expression data and is available for

free at <http://www.informatics.jax.org/mgihome/GXD/GEN/index.shtml>. All data submitted to GXD are given accession IDs that can be referenced in publications and grant applications.

USER SUPPORT

GXD provides users with extensive on-line documentation available by clicking on the question marks available on many pages. In addition, user support staff can be contacted via the 'Help' or 'Contact Us' links on the website, by email (mgi-help@informatics.jax.org) or by telephone (207-288-6445).

CITING GXD

The following citation format is suggested when referring to data from GXD: These data were retrieved from the GXD, and MGI, The Jackson Laboratory, Bar Harbor, Maine, USA (URL: <http://www.informatics.jax.org>). [Type in date (month, year) when you retrieved the data cited.] To reference the database itself, please cite this article.

ACKNOWLEDGEMENTS

We would like to thank our colleagues from the other MGI projects for their contributions to the GXD project and to the larger integrated MGI resource.

FUNDING

National Institutes of Health, Eunice Kennedy Shriver National Institute of Child Health and Human Development grant HD033745. Funding for open access charge: National Institutes of Health grant HD033745.

Conflict of interest statement. None declared.

REFERENCES

- Bult,C.J., Kadin,J.A., Richardson,J.E., Blake,J.A., Eppig,J.T. and the Mouse Genome Database Group. (2010) The mouse genome database: enhancements and updates. *Nucleic Acids Res.*, **38**, D586–D592.
- Gene Ontology Consortium. (2010) The Gene Ontology in 2010: extensions and refinements. *Nucleic Acids Res.*, **38**, D331–D335.
- Begley,D.A., Krupke,D.M., Vincent,M.J., Sundberg,J.P., Bult,C.J. and Eppig,J.T. (2007) Mouse tumor biology database (MTB): status update and future directions. *Nucleic Acids Res.*, **35**, D638–D642.
- Esvikov,A.V., Dolan,M.E., Genrich,M.P., Patek,E. and Bult,C.J. (2009) MouseCyc: a curated biochemical pathways database for the laboratory mouse. *Genome Biol.*, **10**, R84.
- Benson,D.A., Karsch-Mizrachi,I., Lipman,D.J., Ostell,J. and Sayers,E.W. (2010) GenBank. *Nucleic Acids Res.*, **38**, D46–D51.
- Hubbard,T.J., Aken,B.L., Ayling,S., Ballester,B., Beal,K., Bragin,E., Brent,S., Chen,Y., Clapham,P., Clarke,L. et al. (2009) Ensembl 2009. *Nucleic Acids Res.*, **37**, D690–D697.
- Rhead,B., Karolchik,D., Kuhn,R.M., Hinrichs,A.S., Zweig,A.S., Fujita,P.A., Diekhans,M., Smith,K.E., Rosenbloom,K.R., Raney,B.J. et al. (2010) The UCSC Genome Browser database: update 2010. *Nucleic Acids Res.*, **38**, D613–D619.
- Wilming,L.G., Gilbert,J.G., Howe,K., Trevanion,S., Hubbard,T. and Harrow,J.L. (2008) The vertebrate genome annotation (Vega) database. *Nucleic Acids Res.*, **36**, D753–D760.
- Wolfsberg,T.G. (2010) Using the NCBI map viewer to browse genomic sequence data. *Curr. Protoc. Bioinformatics*, Chapter 1, Unit 1.5.1–1.5.25.
- Maglott,D., Ostell,J., Pruitt,K.D. and Tatusova,T. (2007) Entrez gene: gene-centered information at NCBI. *Nucleic Acids Res.*, **35**, D26–D31.
- UniProt Consortium. (2010) The Universal Protein Resource (UniProt) in 2010. *Nucleic Acids Res.*, **38**, D142–D148.
- Hunter,S., Apweiler,R., Attwood,T.K., Bairoch,A., Bateman,A., Binns,D., Bork,P., Das,U., Daugherty,L., Duquenne,L. et al. (2009) InterPro: the integrative protein signature database. *Nucleic Acids Res.*, **37**, D211–D215.
- Amberger,J., Bocchini,C.A., Scott,A.F. and Hamosh,A. (2009) McKusick's Online Mendelian Inheritance in Man (OMIM). *Nucleic Acids Res.*, **37**, D793–D796.
- International Mouse Knockout Consortium, Collins,F.S., Rossant,J. and Wurst,W. (2007) A mouse for all reasons. *Cell*, **128**, 9–13.
- Boon,K., Osorio,E.C., Greenhut,S.F., Schaefer,C.F., Shoemaker,J., Polya,K., Morin,P.J., Buetow,K.H., Strausberg,R.L., De Souza,S.J. et al. (2002) An anatomy of normal and malignant gene expression. *Proc. Natl Acad. Sci. USA*, **99**, 11287–11292.
- Divina,P. and Forejt,J. (2004) The Mouse SAGE Site: database of public mouse SAGE libraries. *Nucleic Acids Res.*, **32**, D482–D483.
- Geschwind,D. (2004) GENSAT: a genomic resource for neuroscience research. *Lancet Neurol.*, **3**, 82.
- Lein,E.S., Hawrylycz,M.J., Ao,N., Ayres,M., Bensinger,A., Bernard,A., Boe,A.F., Boguski,M.S., Brockway,K.S., Byrnes,E.J. et al. (2007) Genome-wide atlas of gene expression in the adult mouse brain. *Nature*, **445**, 168–176.
- Magdaleno,S., Jensen,P., Brumwell,C.L., Seal,A., Lehman,K., Asbury,A., Cheung,T., Cornelius,T., Batten,D.M., Eden,C. et al. (2006) BGEM: an *in situ* hybridization database of gene expression in the embryonic and adult mouse nervous system. *PLoS Biol.*, **4**, e86.
- McMahon,A.P., Aronow,B.J., Davidson,D.R., Davies,J.A., Gaido,K.W., Grimmond,S., Lessard,J.L., Little,M.H., Potter,S.S. and Wilder,E.L. and the GUDMAP project. (2008) GUDMAP: the genitourinary developmental molecular anatomy project. *J. Am. Soc. Nephrol.*, **19**, 667–671.
- Richardson,L., Venkataraman,S., Stevenson,P., Yang,Y., Burton,N., Rao,J., Fisher,M., Baldock,R.A., Davidson,D.R. and Christiansen,J.H. (2010) EMAGE mouse embryo spatial gene expression database: 2010 update. *Nucleic Acids Res.*, **38**, D703–D709.
- Visel,A., Thaller,C. and Eichele,G. (2004) GenePaint.org: an atlas of gene expression patterns in the mouse embryo. *Nucleic Acids Res.*, **32**, D552–D556.
- Smith,C.M., Finger,J.H., Hayamizu,T.F., McCright,I.J., Eppig,J.T., Kadin,J.A., Richardson,J.E. and Ringwald,M. (2007) The mouse Gene Expression Database (GXD): 2007 update. *Nucleic Acids Res.*, **35**, D618–D623.
- Hill,D.P., Begley,D.A., Finger,J.H., Hayamizu,T.F., McCright,I.J., Smith,C.M., Beal,J.S., Corbani,L.E., Blake,J.A., Eppig,J.T. et al. (2004) The mouse gene expression database (GXD): updates and enhancements. *Nucleic Acids Res.*, **32**, D568–D571.
- Ringwald,M., Eppig,J.T., Begley,D.A., Corradi,J.P., McCright,I.J., Hayamizu,T.F., Hill,D.P., Kadin,J.A. and Richardson,J.E. (2001) The Mouse Gene Expression Database (GXD). *Nucleic Acids Res.*, **29**, 98–101.
- Ringwald,M., Eppig,J.T., Kadin,J.A. and Richardson,J.E. (2000) GXD: a gene expression database for the laboratory mouse: current status and recent enhancements. The Gene Expression Database group. *Nucleic Acids Res.*, **28**, 115–119.
- Ringwald,M., Mangan,M.E., Eppig,J.T., Kadin,J.A. and Richardson,J.E. (1999) GXD: a gene expression database for the laboratory mouse. The Gene Expression Database Group. *Nucleic Acids Res.*, **27**, 106–112.

28. Bard,J.B.L., Kaufman,M.H., Dubreuil,C., Brune,R.M., Burger,A., Baldock,R.A. and Davidson,D.R. (1998) An internet-accessible database of mouse developmental anatomy based on a systematic nomenclature. *Mech. Dev.*, **74**, 111–120.
29. Hayamizu,T.F., Mangan,M., Corradi,J.P., Kadin,J.A. and Ringwald,M. (2005) The adult mouse anatomical dictionary: a tool for annotating and integrating data. *Genome Biol.*, **6**, R29.
30. Guo,G., Huss,M., Tong,G.Q., Wang,C., LiSun,L., Clarke,N.D. and Robson,P. (2010) Resolution of cell fate decisions revealed by single-cell gene expression analysis from zygote to blastocyst. *Dev. Cell*, **18**, 675–685.
31. Tamplin,O.J., Kinzel,D., Cox,B.J., Bell,C.E., Rossant,J. and Lickert,H. (2008) Microarray analysis of Foxa2 mutant mouse embryos reveals novel gene expression and inductive roles for the gastrula organizer and its derivatives. *BMC Genomics*, **9**, 511.
32. Shimogori,T., Lee,D.A., Miranda-Angulo,A., Yang,Y., Wang,H., Jiang,L., Yoshida,A.C., Kataoka,A., Mashiko,H., Avetisyan,M. et al. (2010) A genomic atlas of mouse hypothalamic development. *Nat. Neurosci.*, **13**, 767–775.
33. Barrett,T., Troup,D.B., Wilhite,S.E., Ledoux,P., Rudnev,D., Evangelista,C., Kim,I.F., Soboleva,A., Tomaszhevsky,M., Marshall,K.A. et al. (2009) NCBI GEO: archive for high-throughput functional genomic data. *Nucleic Acids Res.*, **37**, D885–D890.
34. Parkinson,H., Kapushesky,M., Kolesnikov,N., Rustici,G., Shojatalab,M., Abeygunawardena,N., Berube,H., Dylag,M., Emam,I., Farne,A. et al. (2009) ArrayExpress update—from an archive of functional genomics experiments to the atlas of gene expression. *Nucleic Acids Res.*, **37**, D868–D872.
35. Ramachandran,S., Ruef,B., Pich,C. and Sprague,J. (2010) Exploring Zebrafish genomic, functional and phenotypic data using ZFIN. *Curr. Protoc. Bioinformatics*, Chapter 1, Unit 1.18.
36. Darnell,D.K., Kaur,S., Stanislaw,S., Davey,S., Konieczka,J.H., Yatskievych,T.A. and Antin,P.B. (2007) GEISHA: an *in situ* hybridization gene expression resource for the chicken embryo. *Cytogenet. Genome Res.*, **117**, 30–35.
37. Bowes,J.B., Snyder,K.A., Segerdell,E., Jarabek,C.J., Azam,K., Zorn,A.M. and Vize,P.D. (2010) Xenbase: gene expression and improved integration. *Nucleic Acids Res.*, **38**, D607–D612.
38. Drysdale,R. and the FlyBase Consortium. (2008) FlyBase: a database for the Drosophila research community. *Methods Mol. Biol.*, **420**, 45–59.
39. Ringwald,M., Baldock,R., Bard,J., Kaufman,M., Eppig,J.T., Richardson,J.E., Nadeau,J.H. and Davidson,D. (1994) A database for mouse development. *Science*, **265**, 2033–2034.